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File: USPT

Sep 28, 1999

US-PAT-NO: 5958672

DOCUMENT-IDENTIFIER: US 5958672 A

TITLE: Protein activity screening of clones having DNA from uncultivated

microorganisms

DATE-ISSUED: September 28, 1999

INVENTOR-INFORMATION:

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CITY

STATE

ZIP CODE

COUNTRY

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CA

US-CL-CURRENT: $\underline{435}/\underline{4}$; $\underline{435}/\underline{183}$, $\underline{435}/\underline{69.1}$, $\underline{536}/\underline{23.1}$, $\underline{536}/\underline{23.2}$

CLAIMS:

What is claimed is:

1. A method for identifying a protein activity of interest comprising:

culturing a gene expression library comprising a pool of expression constructs, each expression construct comprising a vector containing one or more cDNA or genomic DNA fragments, wherein the cDNA or genomic DNA fragments in the pool of expression constructs are derived from a plurality of species of donor organisms, and wherein the cDNA or genomic DNA fragments are each operably-associated with one or more regulatory regions that drives expression of genes encoded by the cDNA or genomic DNA fragments in an appropriate host organism; and

detecting the protein activity encoded by the cDNA or genomic DNA fragments.

- 2. The method of claim 1, wherein the protein activity is an enzymatic activity.
- 3. The method of claim 2, wherein the enzymatic activity is selected from the group consisting of oxidoreductase, transferase, hydrolase, lyase, isomerase, and ligase activity.
- 4. The method of claim 1, wherein the donor organisms are microorganisms.
- 5. The method of claim 4, wherein the microorganisms are derived from an environmental sample.
- 6. The method of claim 4, wherein the microorganisms are a mixed population of uncultured organisms.
- 7. The method of claim 1, wherein the DNA fragment comprises one or more operons, or portions thereof.
- 8. The method of claim 7, wherein the operon or portions thereof encodes a complete or partial metabolic pathway.
- 9. A method for identifying a protein activity of interest comprising:

culturing a gene expression library, comprising a pool of expression constructs,

each expression construct comprising a vector containing one or more cDNA or genomic DNA fragments, wherein the cDNA or genomic DNA fragments in the pool of expression constructs are derived from a plurality of species of donor microorganisms, and wherein the cDNA or genomic DNA fragments are each operably-associated with one or more regulatory regions that drives expression of genes encoded by the cDNA or genomic DNA fragments in an appropriate host organism; and

detecting the protein activity encoded by the cDNA or genomic DNA fragments.

- 10. The method of claim 9, wherein the protein activity is an enzymatic activity.
- 11. The method of claim 10, wherein the enzymatic activity is selected from the group consisting of oxidoreductase, transferase, hydrolase, lyase, isomerase, and ligase activity.
- 12. The method of claim 9, wherein the microorganisms are derived from an environmental sample.
- 13. The method of claim 9, wherein the microorganisms are a mixed population of uncultured organisms.
- 14. The method of claim 9, wherein the DNA fragment comprises one or more operons, or portions thereof.
- 15. The method of claim 14, wherein the operon or portions thereof encodes a complete or partial metabolic pathway.

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